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Sequence Listing was accepted.
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Reviewer: Durreshwar Anjum
Timestamp: [year=2008; month=2; day=19; hr=12; min=44; sec=17; ms=821;]
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Validated By CRFValidator v 1.0.3

Application No: 10576439 Version No: 1.0

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Output Set:

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Finished: 2008-02-08 16:44:25.156
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 499 ms
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Total Errors: 0
No. of SeqIDs Defined: 7
Actual SeqID Count: 7

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SEQUENCE LISTING

<110> Theratechnologies Inc.
Lussier, Bruno
Vachon, Luc
Allas, Soraya
Abribat, Thierry

<120> Selection and treatment of patients suffering from wasting

<130> 09555.0151USWO

<140> 10576439
<141> 2008-02-08

<150> PCT/CA2004/001843
<151> 2004-10-20

<150> 60/512,198
<151> 2003-10-20

<160> 7

<170> PatentIn version 3.3

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<223> Xaa = amino acid sequence of 1 up to 15 residues or is a bond

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Xaa Xaa Asp Ala Ile Phe Tyr Xaa Ser Tyr Arg Lys Xaa Leu Xaa Gln
1   5   10   15

Leu Xaa Ala Arg Lys Leu Leu Xaa Xaa Ile Xaa Xaa Arg Xaa
            20   25   30

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<213> Homo sapiens

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<222> (44)..(44)
<223> Leu residue is capped with an unsubstituted amide moiety

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Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln

1 5 10 15

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg Gln Gln Gly
20 25 30

Glu Ser Asn Gln Glu Arg Gly Ala Arg Ala Arg Leu
35 40

<210> 3
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<212> PRT
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<220>
<223> Amino acid sequence of human GRF

<400> 3

Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln
1 5 10 15

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg Gln Gln Gly
20 25 30

Glu Ser Asn Gln Glu Arg Gly Ala Arg Ala Arg Leu
35 40

<210> 4
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Tyr Ala Asp Ala Ile Phe Thr Asn Ser Tyr Arg Lys Val Leu Gly Gln
1 5 10 15

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg
20 25

<210> 5
<211> 29
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<213> Artificial sequence

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<223> Amino acid sequence of minimum active core of human GRF

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Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg
20 25

<210> 6
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<223> Amino acid sequence corresponding to positions 30 to 44 of human GRF

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Gln Gln Gly Glu Ser Asn Gln Glu Arg Gly Ala Arg Ala Arg Leu
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1 5 10 15

Leu Ser Ala Arg Lys Leu Leu Gln Asp Ile Met Ser Arg Gln Gln Gly
20 25 30

Glu Ser Asn Gln Glu Arg Gly Ala Arg Ala Arg Leu
35 40